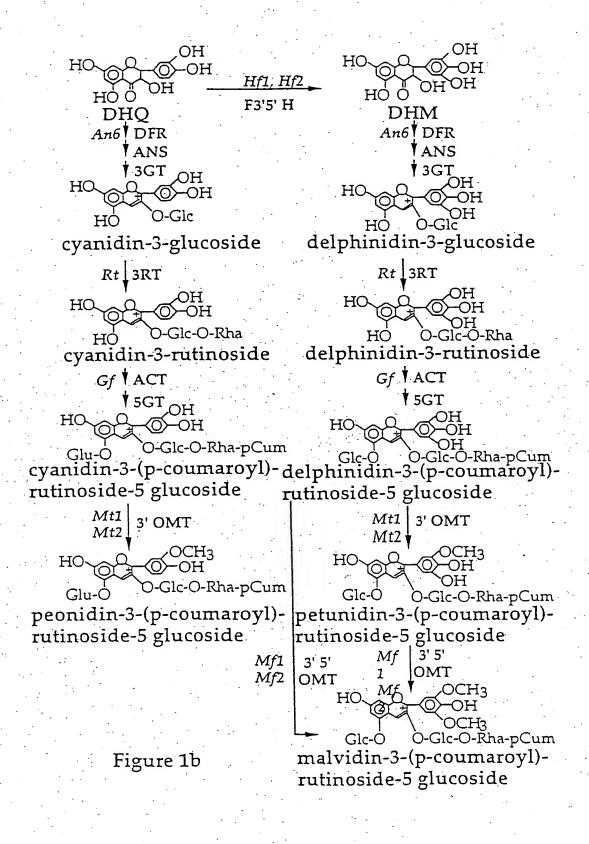


Figure 1a



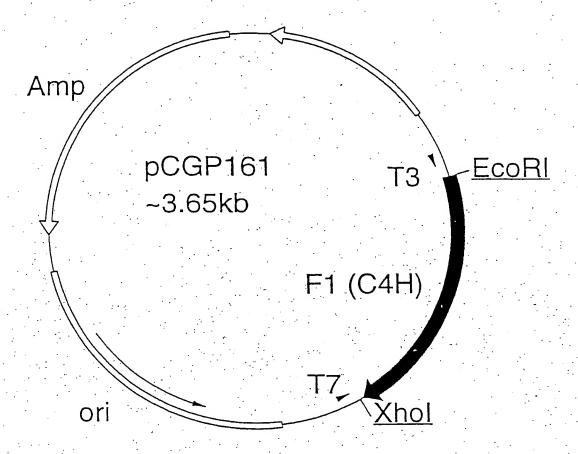


Figure 2

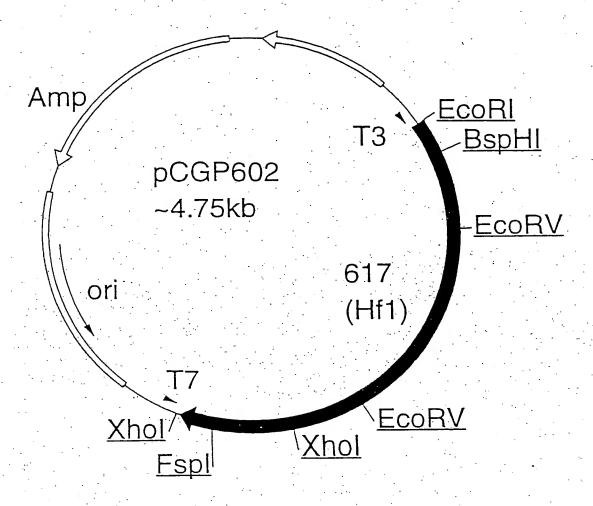


Figure 3

CT/AU97/00124

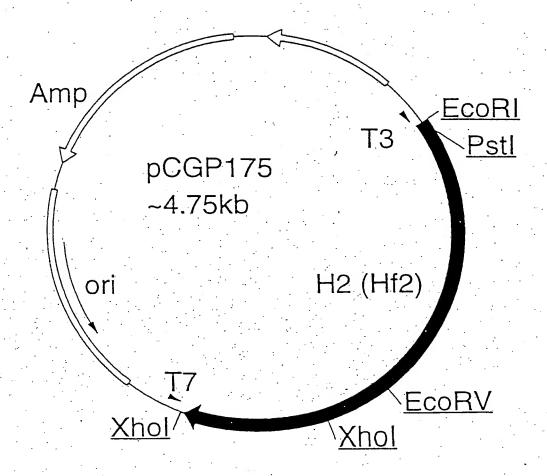


Figure 4

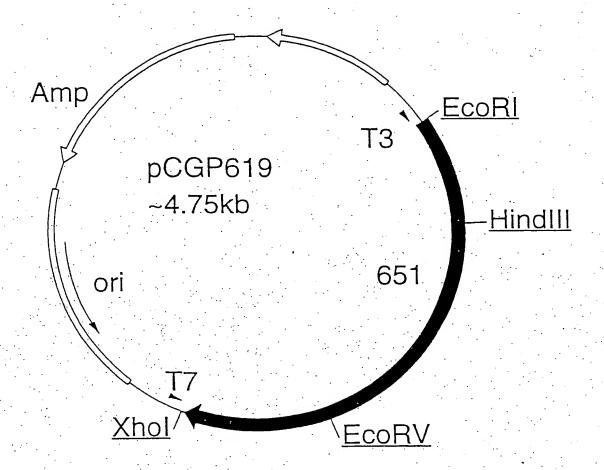


Figure 5



Figure 6

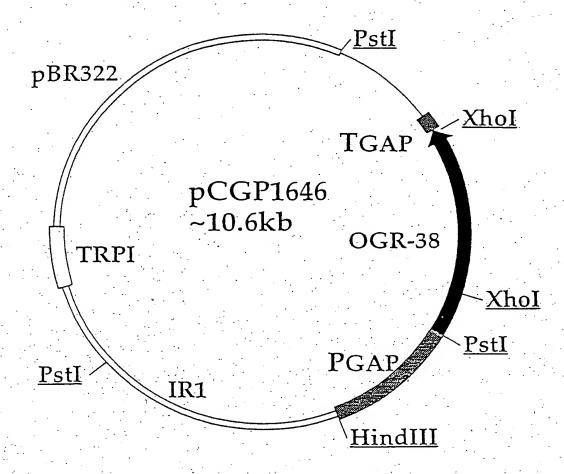


Figure 7

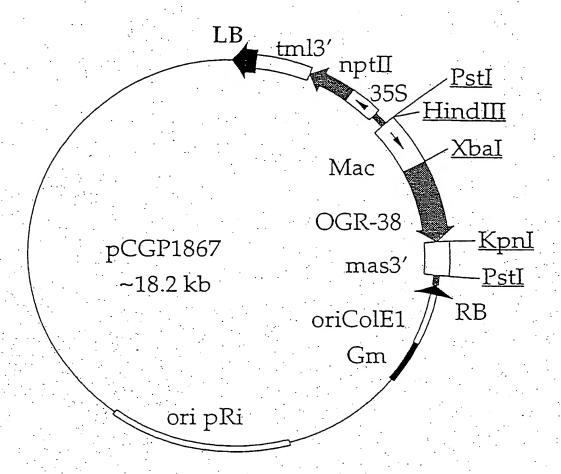


Figure 8

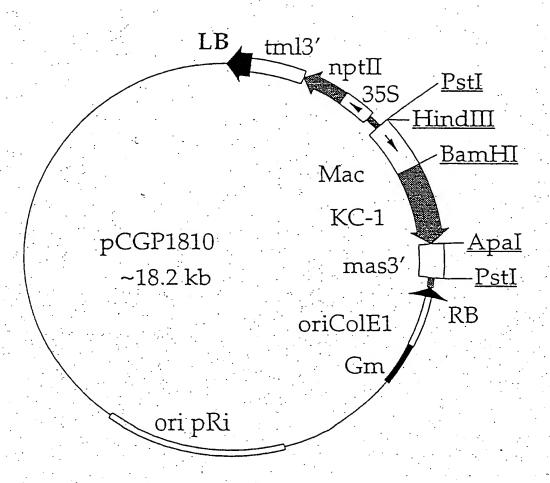


Figure 9

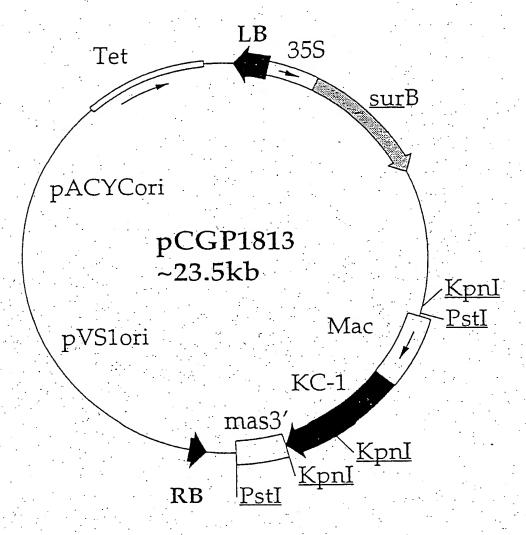


Figure 10

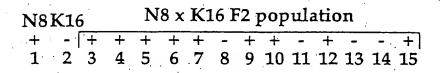




Figure 11

CYANIDIN + CYANIDIN - #1 #3 #4 #5 #8 #6 #11 #12 #13L

**→ → → → 1.8** kb

Figure 12

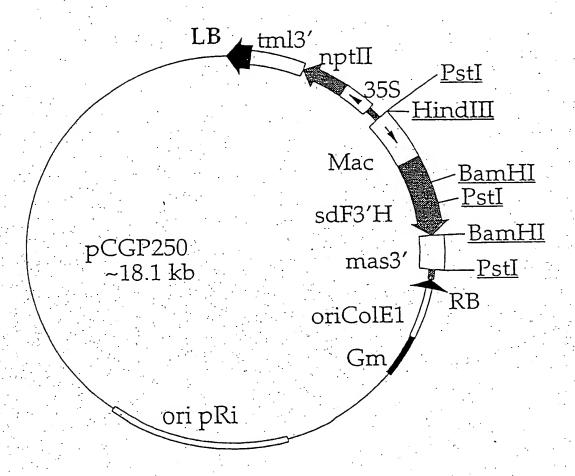


Figure 13

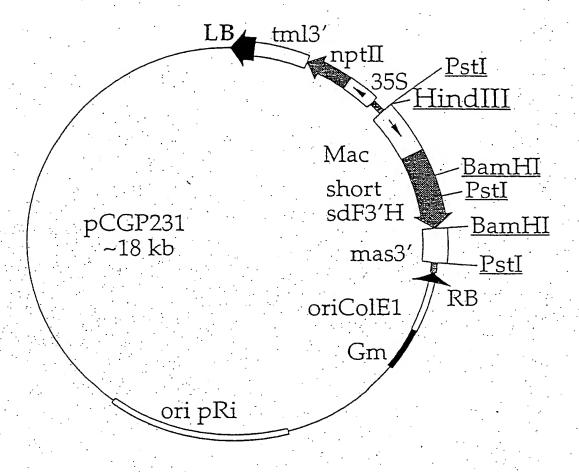


Figure 14

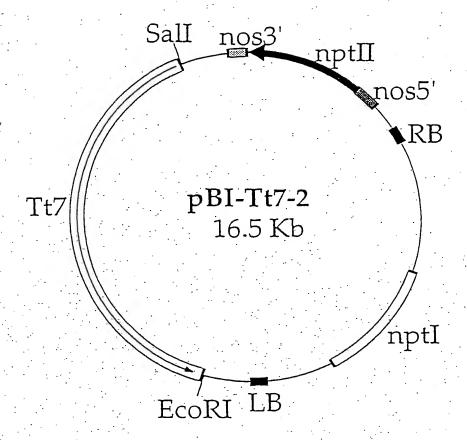


Figure 15

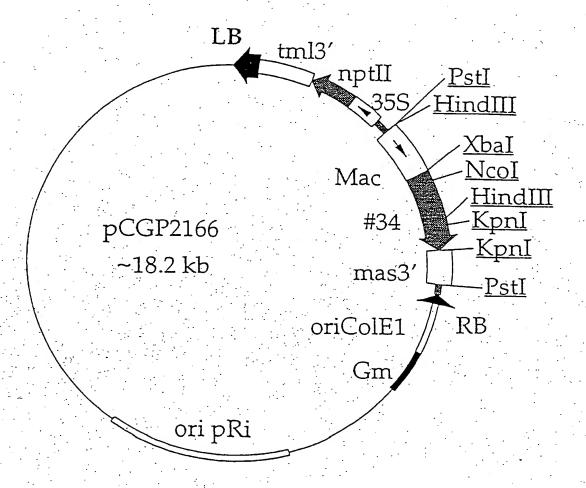


Figure 16

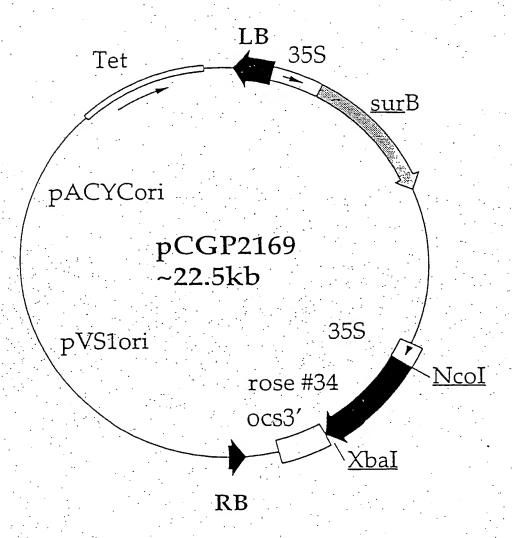


Figure 17

PCT/AU97/00124

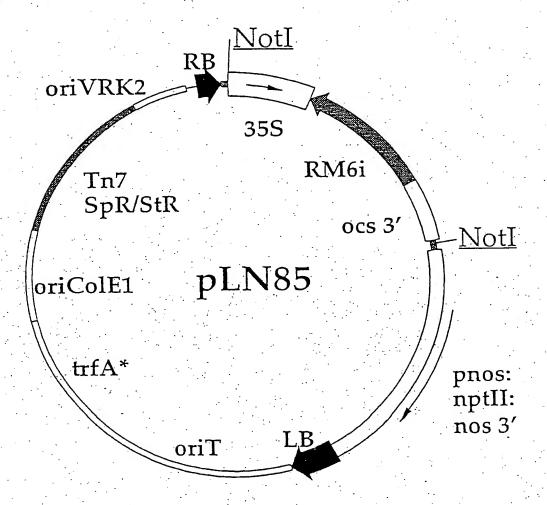


Figure 18

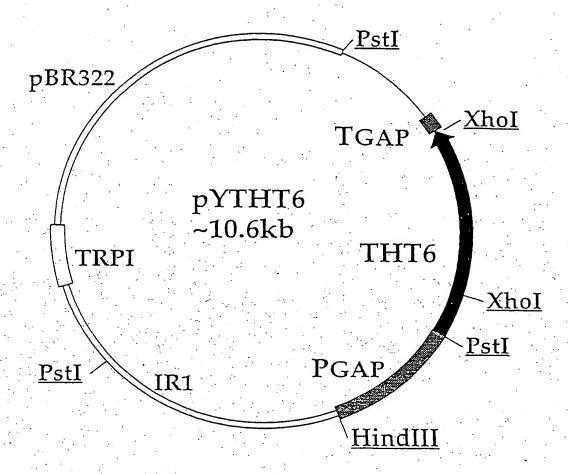


Figure 19

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D
                      mfli|V|vitflfav|FE|frlE|
E
                       lafvfyaLilgsvLyvfL
F
                       l m i I s t l L g f l l Y h s l r L
G
                       1 t 1 I f c t L v f a i F L y f 1 I
H
I
J
                     mdyvn[]llgLfftwF:LvngL
          ffrkRyplpLPPGPKPWPIIGNLVHLG
           ---hqkpLPPGPRPWPIVGNIPHMG
     1 1 y s f 1 n K k v k p L P P G P K P W P I V C N L P H L C
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      fsgksqr-hslpLPPGPKPWPVVCNLPHLC
     nls---sRksarLPPCPtPWPIVGNLPHLG
     1 1 f s g q'g R - - r 1 L P P G P R P W P L V G N L P H L G
G
         -- v k q R y p l p L P P C P K P M P V L C N L P H L C
                                    PILGNIPHEG
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   37 Q a PR Q G L X X L X Q K Y G P L L Y M R L G Y V D V V X
В
      PK FHQSMAATARV hGFLI HLKMGFVhVVVA
C
   61
      t X PHRTLS AMV t tY GPILELRLGFVDVVVA
D
      pfPHHSIAeLAKkhGPLMHLRLGYVDVVX
E
      PIPHHALAALAKKYGPLMHLRLGCVDVVVA
F
      PK-P-Has MAeLARaY C-PLMHLKMG FVhVVVA
G
    48 · k K PH Q S I A A M A e r Y G P L M H L R L G F V D V V A
H
      S K P H Q T L A E M A K L Y G P L M H L K f G 1 k D a V V A
      nHPHKSLAqLAKihGPIMNLKLGqLntVVi
      ASASVA a QFLK THD ANPSSRPPNSG A e HMA
      AS ASS A LOFIK HDINFS SEPPNSGAKHIA
В
      S S A S V A e k P L K V H D A N F S S R P P N S G A K H V A
      ASKS VAEQ PLKIHDAN PASR PPN SCAKHMA
D
      A S A S V A a Q F L K t H D A N F S S R P P N S G A K H L A
E
      A S A S V A a Q P L K V H D A N F A S R P P N S G A K H V A
       S S. A S a A | e | Q C L R | v | H D A N F 1 S R P P N S G A K H V A
 G
      ASAAVAAQPLKVHDSNFSNRPPNSCAEHIA
                                                    107
    78
        SASVA e O P L K K H D V N F S n R P P N S G A K H I A
            V A r E v L Q k Q D l T P S n R f v p d v v H v r n
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YNY Q DLVFAPTCPKWRMLRKICSLHMFSSK
C
      YNYODLV PAPYGPRWRMLRKICALHLPSAK
                                                   150
  121
      Y NYO DLV PAPYCh RWRLLR KISSV HLFS AR
D
                                                   140
  111
      THY QOOLLY FREE CPRING MERKISSVHLFSGK
E
  109
      YN Y Q DLV FAPY GPRWRLLRKICS V H L P S A K
   109
      YNY EDILVP P PYGPKWR L LRKICA GHIPSVK
G
  112
      YNYODL V PAPIC PRWRM LRKIT SV HLPSAK
Η
  108
                                                   137
      YNYODLVFAPYGPRWRLLRKICSVHLFSS
                                                   100
   71
      hsdfsV:VwlPvNsR-WKtLRK-ImnssIFSgn
     ALDDPRHVRQD---EVktLTRALASAGqkP
  127 ALDDOFRIVER OF
                      - - - E"V S I L V n A I A k A G t. k P
В
                      - - E V G I L T R V L A d A C e t P
С
     AL n D P L E V R Q D
  151
                      --- EVgtLTReLvrvGtkP
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D
  I41
                      - - - R V S V L A H A L A n S G s k v
E
      A L D D 1 K H V R Q E
  139.
                         - EVAVETRVLISAGNSP
· F · 139
      ALD DOP R HOVEROME
                                           -]c|kr
G 142
                         - EVALLSRALA
                                                   166
      A M D DEP R R VER E E
                           EVALLTRSLASAGKEP
H
      ALDDFCHVROE
                                                   164
  . 138
      ALD DEP CH VER hE
                                                   127
I.. 101
                           EICILIRAIAS gGhaP
                             r I Trans I As A Gen P
  \cdot , 1
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      VOLCOLLNVCTTNALSRVHLCKRV1CdCtC
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C
      VNLCQLVNMCVVNALGREMIGRRLFG---
D
  168
      V N LaO L L N L C T V N A L G R V M V G R R V F G d G s G
E
  166
      VOLCQLINVCATNALARVMLCRRVFC - - - d
F
   166
      VPICOMEN VCATNACESRVMMGRRVVChadC 196
G
   167
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H
   .165
      VNLGK:LLGVCTTNALARVELGRRVFE-GdG
   128
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   169.
A 198 d v D P Q A a E F K S M V V E M M V V A G V F N I G D F I P
В
   184
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                                                    213
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                                                    233
   207
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                                                    224
D
   195
      g d D P K A D E P K S M V V E M M V L A G V P N I G D
                                                    .225
      g i Dr s A n e P K d M V V E L M V L A G e P N L
F
   197 tnDaKAEBPKAHVVELMVLSGVFNIGD
G
      k s D P X A E E P X S M V e E M M V L A G S P N I G D
                                                    224
                                                    186
      gen P H A D E F K S M V V E I M V L A G a F N L G D
   45 g v D P Q A D E P K S M V V E I M V L A G a P N L G D P I P
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 K 195. pfsdsAkerKelvwnIMVeAckpNLVDYfp
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D
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E
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F
     VIDII f D.L.Q.CITKKWXXXLHVRPDSPLSKIVE
G
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     VLgWfDVQGIVgXXXLEARPDAFLncILE
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B 244
     DHKINss-detkgHs-DLInmLISLKdadd
C
 264
     E H e m Ng - - - q d q K h t D M L S T L I S L X g t d l
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  .255
E 256
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G
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H
 255
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 : 217
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     EHKSNgsr - - gakQhvDL SmLISLQdnni
     DRlkern - - 1 rdnani DV LdAL Ln Isqenp
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  283 DNdg - - GKLTDTEIKALLINLPVAGTDTSS
     DG--eCCKfSaTETXALLELGUPTACTDTSS
В
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     ae---CCRLTDVEIXALLINNLPAAGIDITIS
 - 292
                                              318
D
  281
     DG - - d GFG S LET D T E I K A L LEN M F T A G T D T S A
                                              -308
        - - e CC K L T D T E I K A L L L N M P T A C T D T S S
E
  281
                                              308
\mathbf{F}
  279
        ---GOOK LOTE I KALLIEN LIPAAGT DISS
                                               305
G
        --- GSRLTDTEIKALLLENLI----DTTS
  286
                                              30%
H
  285
     DmdceeGKLTDTEIKALLLKLFTAGTDTSS
                                              314
Ι
  245
     D C g d e C t K L T D T E I K A L L L N L F i A G T D T S S
J.
     DG-esGaKLTDTEIKALLENTFTAGTDTSS
            --- Id r N Q I e q L c L d L F A A G T D T T S
A
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                                               340
В
  297
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C
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D
  309
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E
     STIVE WAI A ELIRHPHM LaRVOKELD dfVGH
F
     S.T V R W A I A E L I R H P Q I L k Q A R E E I D a V V G Q
· G
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C
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                                                 368
D
  339
      dary TesdiphipricoxvixerPalesTP
E
                                                 368
      dRIVTEIDLSQLCTIQALVKEVFRLHPSTP
                                                 365
     n R L V S R T D L n O L P Y L O A V V K E T P R L H P P T P
                                                 368
G
     n Q L V T E S D L t d L P F L Q A I V K E T F R L H P S T P
                                                 374
     n R f V T E S D L p Q L t F L Q A V I K E T F R L H P S T P
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  . 192
      g K q V e E A D V g r L P Y L r C I V K E T l R I H P
K
  337
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                                                 400
A
                                                 386
  357
     LES LEPERVAN ZES CE INGY h I PK NTTLL VNVWA
B.
     LSLPRIAMESCEVNCYIIPKCSTLL
C
  379
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                                                 398
D
  369
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                                                 198
E
  369
                                                 395
     LES LERRISSESCE V d CYYLP K CSTLL V N V W A
F.
  366
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G
  369
      ISLPRMGAQGCEINCY fIPKCATLUVNVWA
H
  . 375
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      LlIPRK veEdvELstYiI:PKdSqVLVNVWA
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      TO REPERVE OF LEGGEKPNVDV
C
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D
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E
   399
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F
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                                                 428
G
   399
      LARDPNVWTNPLEFNPhRFLPGGEKPNVDI
      TARDOPRAMANDOLDENPLECERPNV
   365
                                                 281
      IARDPSIWADPLEFRPARFLPGGEKPNVDV
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K
  397
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   431
      KG M D PELIPP GAGRRICAGL SLGLRMV
 В
   417
 C
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                                                 - 468
             ELIPPGAGRRICAGLSLGLRE I Q E L
                                                  458
 D
      RGNDFEVIPFGAGRRICAGMSLGLRMVH
                                                  458
 Ê
   429
      RGNDFEVII PFGAGRRICAGASLGLRMVQ
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 F.
   426
                                                  458
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 G
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                                                  464
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 Η
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   395
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   282
      R G T D F E L I P F G A G R R I C D G L D L a M R M I D V
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С	469	TATL OHAPO f D LA d G Q 1 PE S L N K E E A Y C L T	498
D	459	TATLEY Q G PODEM E LA G G V t PE K LEN H P E S Y G L T	488
E	459	TATLVHAFRWALADGLEAEKLHMDEAYCLT	488
F	456	i A T L V Q T F D W E L A N C L e P E M L N M R R A Y C L T	485
G	459.	TASLIRAPDID LANGLIA ON LN MERAY GLT	488
H	465	VATILUE A Z D M D L V N G G S V Z C L N M E B A Y C L T	494
Ţ	425	TWASLVESTED WALL de L KPEKLdke BEGTOCLT	454
٠	312	TATLUTES FIND LING MS PC KLCHERNYCLT	341
K	453	1 gs L Lns Fn Wk LyG G I a Pk d L d M g Ek F G I T	482
1.	733	Talandarian Mymako a latik amamarki e T.L.	<del>7</del> 02
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Α.	401	TOOR ADDRESS VVVV PRRRESSO SO VI	513
A	491 477	LOORAD PELVVHIPRPRLeaQaYig	512
В	477	LORK V PL M VIH PtrRLSaRVYnsgf	500
В	477 499	LORKVPLMVHPtrRLSaRVYnsgt LORAdPLVVHPKPR	500 512
B C D	477 499 489	LORK V PL M V H P t r R L S a R V Y n s g f LOR A d P L V V H P K P R LOCK A V P L V V H P K P R L A P n V Y g 1 g s g	500
B C D E	477 499	LORK V PL M V H P t r R L S a R V Y n s g f L Q R A d P L V V H P K P R L A P n V Y g 1 g s g L O R A a P L M V H P R t R L A P Q a Y k t s s s	500 512
B C D	477 499 489	LORK V PL M V H P L T R L S A R V Y N S G f L Q R A d P L V V H P K P R L O R A V P L V V H P K P R L A P N V Y G 1 G S G L O R A A F L M V R P R L R L A P Q A Y K t S S S	500 512 513
B C D E	477 499 489 489	LORK V PL M V H P t r R L S a R V Y n s g f L Q R A d P L V V H P K P R L A P n V Y g 1 g s g L O R A a P L M V H P R t R L A P Q a Y k t s s s	500 512 513 512
B C D E F	477 499 489 489 486	LORA DEL MVHP KPRLSARVY nsgf LORA DEL VVHP KPR LOCRA V PLVVHP KPRLAP NVY glgsg LORA A EL MV FPRLRLAP QAY ktsss LORA A FLMV HPRPRLAPHVY esi LORA A PL VHPRPRPRLA HVY	500 512 513 512 508 508
B C D E F G	477 499 489 489 486 486	LORA VPL M VH P t r R L S a R V Y n s g f LORA VPL V V H P K P R LORA VPL V V H P K P R L A P n V Y g 1 g s g LORA a P L M V H P R T R L A P H V Y e s i LORA a P L M V H P R P R L A L H V Y LORA e P L L V H P R P R L A L H V Y LORA V P L M L H P R P R L Q P H L Y t 1 n	500 512 513 512 508 508 517
B C D E F G H I	477 499 489 489 486 489 495 455	LORK V PL M V H P L T R L S a R V Y N s g f L Q R A d P L V V H P K P R L Q R A v P L V V H P K P R L A P N V Y g 1 g s g L Q R A a P L M V H P K P R L A P H V Y e s i L Q R A a P L M V H P K P R L A L H V Y L Q R A e P L L V H P R P R L A L H V Y L Q R A v P L M L H P K P R L Q P H L Y L 1 n L Q R A s P L I V H P K P R L S a Q V Y c m	500 512 513 512 508 508 517 476
B C D E F G H I	477 499 489 489 486 489 495 455 342	LORA VPL M VH P t r R L S a R V Y n s g f LORA VPL V V H P K P R LORA VPL V V H P K P R L A P n V Y g 1 g s g LORA a P L M V H P R T R L A P H V Y e s i LORA a P L M V H P R P R L A L H V Y LORA e P L L V H P R P R L A L H V Y LORA V P L M L H P R P R L Q P H L Y t 1 n	500 512 513 512 508 508 517